

OIPE

1642

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/848,852ADATE: 12/20/2001  
TIME: 11:52:09Input Set : N:\Crf3\RULE60\09848852.raw  
Output Set: N:\CRF3\12202001\I848852A.rawRECEIVED  
JAN 18 2002  
TECH CENTER 1600/2900

ENTERED

## SEQUENCE LISTING

## 3 (1) GENERAL INFORMATION:

5 (i) APPLICANT: Hillman, Jennifer L.  
6 Tang, Y. Tom  
7 Corley, Neil C.  
8 Guegler, Karl J.  
9 Yue, Henry  
10 Patterson, Chandra

12 (ii) TITLE OF INVENTION: PROTEINS ASSOCIATED WITH EGF SIGNALING

14 (iii) NUMBER OF SEQUENCES: 5

16 (iv) CORRESPONDENCE ADDRESS:

17 (A) ADDRESSEE: Incyte Pharmaceuticals, Inc.  
18 (B) STREET: 3174 Porter Dr.  
19 (C) CITY: Palo Alto  
20 (D) STATE: CA  
21 (E) COUNTRY: USA  
22 (F) ZIP: 94304

24 (v) COMPUTER READABLE FORM:

25 (A) MEDIUM TYPE: Diskette  
26 (B) COMPUTER: IBM Compatible  
27 (C) OPERATING SYSTEM: DOS  
28 (D) SOFTWARE: FastSEQ for Windows Version 2.0

30 (vi) CURRENT APPLICATION DATA:

C--> 31 (A) APPLICATION NUMBER: US/09/848,852A  
C--> 32 (B) FILING DATE: 07-May-2001

34 (vii) PRIOR APPLICATION DATA:

35 (A) APPLICATION NUMBER: 09/069,725  
36 (B) FILING DATE:

40 (viii) ATTORNEY/AGENT INFORMATION:

41 (A) NAME: Billings, Lucy J.  
42 (B) REGISTRATION NUMBER: 36,749  
43 (C) REFERENCE/DOCKET NUMBER: PF-0515 US

45 (ix) TELECOMMUNICATION INFORMATION:

46 (A) TELEPHONE: 650-855-0555  
47 (B) TELEFAX: 650-845-4166

50 (2) INFORMATION FOR SEQ ID NO: 1:

52 (i) SEQUENCE CHARACTERISTICS:

53 (A) LENGTH: 298 amino acids  
54 (B) TYPE: amino acid  
55 (C) STRANDEDNESS: single  
56 (D) TOPOLOGY: linear

58 (vii) IMMEDIATE SOURCE:

59 (A) LIBRARY: COLNNOT16  
60 (B) CLONE: 1281694

62 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

64 Met Asn Asp Ser Leu Arg Thr Asp Val Phe Val Arg Phe Gln Pro Glu

65 1

5

10

15

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/848,852A

DATE: 12/20/2001  
TIME: 11:52:09

Input Set : N:\Crf3\RULE60\09848852.raw  
Output Set: N:\CRF3\12202001\I848852A.raw

66 Ser Ile Ala Cys Ala Cys Ile Tyr Leu Ala Ala Arg Thr Leu Glu Ile  
67 20 25 30  
68 Pro Leu Pro Asn Arg Pro His Trp Phe Leu Leu Phe Gly Ala Thr Glu  
69 35 40 45  
70 Glu Glu Ile Gln Glu Ile Cys Leu Lys Ile Leu Gln Leu Tyr Ala Arg  
71 50 55 60  
72 Lys Lys Val Asp Leu Thr His Leu Glu Gly Glu Val Glu Lys Arg Lys  
73 65 70 75 80  
74 His Ala Ile Glu Glu Ala Lys Ala Gln Ala Arg Gly Leu Leu Pro Gly  
75 85 90 95  
76 Gly Thr Gln Val Leu Asp Gly Thr Ser Gly Phe Ser Pro Ala Pro Lys  
77 100 105 110  
78 Leu Val Glu Ser Pro Lys Glu Gly Lys Gly Ser Lys Pro Ser Pro Leu  
79 115 120 125  
80 Ser Val Lys Asn Thr Lys Arg Arg Leu Glu Gly Ala Lys Lys Ala Lys  
81 130 135 140  
82 Ala Asp Ser Pro Val Asn Gly Leu Pro Lys Gly Arg Glu Ser Arg Ser  
83 145 150 155 160  
84 Arg Ser Arg Ser Arg Glu Gln Ser Tyr Ser Arg Ser Pro Ser Arg Ser  
85 165 170 175  
86 Ala Ser Pro Lys Arg Arg Lys Ser Asp Ser Gly Ser Thr Ser Gly Gly  
87 180 185 190  
88 Ser Lys Ser Gln Ser Arg Ser Arg Ser Arg Ser Asp Ser Pro Pro Arg  
89 195 200 205  
90 Gln Ala Pro Arg Ser Ala Pro Tyr Lys Gly Ser Glu Ile Arg Gly Ser  
91 210 215 220  
92 Arg Lys Ser Lys Asp Cys Lys Tyr Pro Gln Lys Pro His Lys Ser Arg  
93 225 230 235 240  
94 Ser Arg Ser Ser Arg Ser Arg Ser Arg Ser Arg Glu Arg Ala Asp  
95 245 250 255  
96 Asn Pro Gly Lys Tyr Lys Lys Ser His Tyr Tyr Arg Asp Gln Arg  
97 260 265 270  
98 Arg Glu Arg Ser Arg Ser Tyr Glu Arg Thr Gly Arg Arg Tyr Glu Arg  
99 275 280 285  
100 Asp His Pro Gly His Ser Arg His Arg Arg  
101 290 295  
103 (2) INFORMATION FOR SEQ ID NO: 2:  
105 (i) SEQUENCE CHARACTERISTICS:  
106 (A) LENGTH: 3464 base pairs  
107 (B) TYPE: nucleic acid  
108 (C) STRANDEDNESS: single  
109 (D) TOPOLOGY: linear  
111 (vii) IMMEDIATE SOURCE:  
112 (A) LIBRARY: COLNNOT16  
113 (B) CLONE: 1281694  
115 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:  
117 TTCTGTGAAGC ACTCCATGG A GCATGTGTCA ATGGCCTGTG TCCACCTGGC TTCCAAGATA 60  
118 GAAGAGGCC CAAGACGCAT ACGGGACGTC ATCAATGTGT TTCACCGCCT TCGACAGCTG 120  
119 AGAGACAAAA AATAATCGTT ATGTACCTTC AGGTGTTAGA GTGTGAGCGT AACCAACACC 180

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/848,852A

DATE: 12/20/2001

TIME: 11:52:09

Input Set : N:\CrF3\RULE60\09848852.raw

Output Set: N:\CRF3\12202001\I848852A.raw

120	TGGTCCAGAC	CTCATGGGTA	GCCTCTGAGG	GTAAGTGA	AAGACTTCTC	CTCTGCTGTC	240
121	CAAGCGCTT	GGTGCAGGGA	CAGCGGCATC	TTCAGCCAAT	CCAGTGCAGG	CTCTCCACCG	300
122	AAGGCTGGCT	CTAGACTGGT	GACCCCTGT	TGAAATGGG	CAGTTGGCAG	CGGCTCTGAT	360
123	GAGCCCGAGA	AGAGGCCTGC	CCTTGGGTGC	GGAGTCTCCC	TCCGCACGAT	GCTCCCACGC	420
124	GTCCAACTTG	CACCCAAGGG	GCTTTCCCT	CTTCCAAGTG	GACTCCTCA	AGGAAGCTGC	480
125	AGCTCGGTCA	GCAGAGAAGG	GGCCTGCCGC	CAGCGCCCTG	GAGGAAGAGG	AAGAGGAACC	540
126	CAAGAGGATG	GCTTGTCTCC	CAGCAGCCAC	ACCGGCTTTG	TGCTCAGCCA	GTTCATTG	600
127	GTTTGCATGT	TTCTCTGCAC	TATGGATTT	GAGCATTAG	ATTTCTTAA	TCAAAAGCGT	660
128	TTTAGTGA	CCAGTAGACA	TTTCTTTCT	GAGGCATCGT	GCTTTGCATG	AGAGCAGGCC	720
129	AAGGTTGAGG	GGAAAAGTAA	AGTTAAAGTC	GGTCTCTTT	CATAGCAACA	CGTATTGTCT	780
130	GACATTCA	CAGCTTTTT	TTTTCTAA	AATTCTGTG	CCTTCCTGTC	CTGTATTAC	840
131	TGTATTTAGA	AAAAGCAGCT	AGAATATTC	TCCATTA	CTTGAGATT	ACAGGACTGT	900
132	CTAGCTCTGA	GTCCTAGCAA	TAGACTCCTT	AGAGGAGTAG	TACGTTTATC	TAGATTTCT	960
133	CTAGATAATG	CAGGCGGAAG	ACCTGGGTT	CCGGGTGGGG	CATTGCAGTT	CTTCCTGTGT	1020
134	TTGGCTTCA	GGAAATTACAT	GAACGACAGC	CTTCGCACCG	ACGTCTCGT	GCGGTTCCAG	1080
135	CCAGAGAGCA	TCGCCTGTGC	CTGCATTAT	CTTGCTGCC	GGACGCTGG	GATCCCTTTG	1140
136	CCCAATCGTC	CCCATTGGTT	TCTTTGTT	GGAGCAACTG	AAGAAGAAAT	TCAGGAAATC	1200
137	TGCTTAAAGA	TCTTGCAGCT	TTATGCTCG	AAAAAGGTTG	ATCTCACACA	CCTGGAGGGT	1260
138	GAAGTGGAAA	AAAGAAAGCA	CGCTATCGAA	GAGGCAAAGG	CCCAAGCCG	GGGCCTGTTG	1320
139	CCTGGGGGCA	CACAGGTGCT	GGATGGTACC	TCGGGGTTCT	CTCCTGCC	CAAGCTGGT	1380
140	GAATCCCCA	AAGAAGGTAA	AGGGAGCAAG	CCTCCCCAC	TGTCTGTGAA	GAACACCAAG	1440
141	AGGAGGCTGG	AGGGCGCCAA	GAAAGCCAAG	GC GGACAGCC	CCGTGAACGG	CTTGCCAAAG	1500
142	GGGCGAGAGA	GTCGGAGTCG	GAGCCGGAGC	CGTGAGCAGA	GCTACTCGAG	GTCCCCATCC	1560
143	CGATCAGCGT	CTCCTAACAGAG	GAGGAAAAGT	GACAGCGGCT	CCACATCTGG	TGGGTCCAAG	1620
144	TCGCAGAGCC	GCTCCCGGG	CAGGAGTGC	TCCCCACCGA	GACAGGCCCC	CCGCAGCGCT	1680
145	CCCTACAAAG	GCTCTGAGAT	TCGGGGCTCC	CGGAAGTCCA	AGGACTGCAA	GTACCCCCAG	1740
146	AAGCCACACA	AGTCTCGGAG	CCGGAGTCT	TCCCGTTCTC	GAAGCAGGTC	ACGGGAGCGG	1800
147	GCGGATAATC	CGGGAAAATA	CAAGAAGAAA	AGTCATTACT	ACAGAGATCA	GCGACGAGAG	1860
148	CGCTCGAGGT	CGTATGAACG	CACAGGCCGT	CGCTATGAGC	GGGACCACCC	TGGGCACAGC	1920
149	AGGCATCGGA	GGTGAGGCCG	GGTTGCAGTG	ACTGGTGGCC	GCAAGCCCT	CCCTGGGGAG	1980
150	TACCTGATGG	CTGCCCTTTG	ACCCCCGGT	GCTGCCCTT	GACCCCGGG	TGTGCTCTCA	2040
151	GCGCAAGTGG	TCCTAGAAC	GGATTCTTT	TGAAATGTC	TGTCGACTGG	ACCTTGGTGG	2100
152	ATTGGAAT	GGAACTGAGG	GACCGGTGAC	ACGTGCTTC	GACCGGTCTG	GGGTGCGGCG	2160
153	CACACCTGGG	CCCCTGCAGG	GCTCAGCTCG	GCAGCAGCTC	TGAGGGCAGC	TCAATGAAA	2220
154	AGTGAATGCA	CACGCCCTG	TTGGCGTGGC	CTGGCATGGC	CTGGTGTAT	CGGCAGCCGC	2280
155	TCTCCACTCC	CCGACTGATA	CTCAATTACG	TGAAGCCAAG	AAAGATGATT	TTTGAACCT	2340
156	TTGCCTATAT	TAGGTTGTAC	TTATGTACAT	ATTTGCA	GTTCACAGG	AGAAAGTGGC	2400
157	CTTAAC	TGCCCTTCTC	TCTCCACGTT	GTAAATAAAC	ATGTGTTAA	TACAAGTAA	2460
158	AGCTATGTAT	GAAAAC	AACTTGAATC	CCGTCA	AAAAC	TTGGAATCC	2520
159	TGACTTTAA	AATGTGAGGG	TATTTGGATC	TGTGTTGAAA	GTCGTATATT	TTTATCTGTG	2580
160	CGGTGCTGAG	TGCAGGCCAC	CAGCTCTAA	ATAGAGGTT	CCTATATGCG	CGTATGACAT	2640
161	GGTGAATAAA	CACAACTCTC	TCCACTCAGG	ACATCCGGAG	CGTTATGGAC	GTGGTAGGTG	2700
162	GTCGTTCTGT	GTGCTTGTGA	AAGTGTCCAG	GCGTGTGAC	AGCCAGTGCG	CCCACTTCCG	2760
163	GGCTCCTTGC	TCCCTGCTGT	ACTGAAGTT	TGGATTTGC	ATCCAATCCT	GTGTGCCTGC	2820
164	CCTTCTGCCG	AAGCTTGTGA	GGGGCCTGAG	TCCCTGCC	ATCAGGATGA	CAGGCTCCTT	2880
165	CCTGCAGGGC	CATAGGAGGG	AAGTTTGG	AAACACAGAAT	GATTCCAAGG	TGCTCTCGTT	2940
166	CCTGAGGGGG	ACTGTTTGT	AACCCATGAC	ATCTGTGGC	GAGAGAGGCA	GCTGGGAGCA	3000
167	GGACACTTGG	AGGGTCACCC	CACGGGGTG	GCACCTGCAC	TCTGAGTGC	CCCCACTGTC	3060
168	ATCAGCTGCC	TCTTACCGTG	GACACAGTT	TGGTTTGGG	GACTAGGGGG	CCCCACTCCT	3120

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/848,852A

DATE: 12/20/2001  
TIME: 11:52:09

Input Set : N:\Crf3\RULE60\09848852.raw  
Output Set: N:\CRF3\12202001\I848852A.raw

169 GGTGGTACCG TTTGGACTTA CTAGGGCAGT GGGACATATA GGCGGGGCT AGTGGGATAA 3180  
170 CGGGGAGTTA CGCCTGATGA CTTTTTGAT GGAATCCTGC ATTAGATAGC TGGTGGGACC 3240  
171 CCCCCCTCAG AATTGGGGAA CTGAGGAGAC TCCAGGGAGG GTGTCTTCC AGGGAGAGCA 3300  
172 GCTATGAGGG GCCCCCTAGC TTCCTGTGCC TGGAAAGTAAG AGAACCAAGTA AAGGGCCATA 3360  
173 CACACCTGTA CCCAAGAGAC CGCTCTCCAT TTGCTTTCTT TTTTTACTAA ATAATTGTAA 3420  
174 AATATTATTA TGACATAAAG AACCATTAA GGCCAAAAAA AAAA 3464

176 (2) INFORMATION FOR SEQ ID NO: 3:

178 (i) SEQUENCE CHARACTERISTICS:  
179 (A) LENGTH: 332 amino acids  
180 (B) TYPE: amino acid  
181 (C) STRANDEDNESS: single  
182 (D) TOPOLOGY: linear

184 (vii) IMMEDIATE SOURCE:

185 (A) LIBRARY: BEPINOT01  
186 (B) CLONE: 2056178

188 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

190 Met Tyr Ser Ala Gln Arg Phe Trp Gly Thr Ile Trp Ala Arg Arg Gly  
191 1 5 10 15  
192 Ala His Leu Ala Pro Pro Asp Ala Ser Ile Leu Ile Ser Asn Val Cys  
193 20 25 30  
194 Ser Ile Gly Asp His Val Ala Gln Glu Leu Phe Gln Gly Ser Asp Leu  
195 35 40 45  
196 Gly Met Ala Glu Glu Ala Glu Arg Pro Gly Glu Lys Ala Gly Gln His  
197 50 55 60  
198 Ser Pro Leu Arg Glu Glu His Val Thr Cys Val Gln Ser Ile Leu Asp  
199 65 70 75 80  
200 Glu Phe Leu Gln Thr Tyr Gly Ser Leu Ile Pro Leu Ser Thr Asp Glu  
201 85 90 95  
202 Val Val Glu Lys Leu Glu Asp Ile Phe Gln Gln Glu Phe Ser Thr Pro  
203 100 105 110  
204 Ser Arg Lys Gly Leu Val Leu Gln Leu Ile Gln Ser Tyr Gln Arg Met  
205 115 120 125  
206 Pro Gly Asn Ala Met Val Arg Gly Phe Arg Val Ala Tyr Lys Arg His  
207 130 135 140  
208 Val Leu Thr Met Asp Asp Leu Gly Thr Leu Tyr Gly Gln Asn Trp Leu  
209 145 150 155 160  
210 Asn Asp Gln Val Met Asn Met Tyr Gly Asp Leu Val Met Asp Thr Val  
211 165 170 175  
212 Pro Glu Lys Val His Phe Phe Asn Ser Phe Phe Tyr Asp Lys Leu Arg  
213 180 185 190  
214 Thr Lys Gly Tyr Asp Gly Val Lys Arg Trp Thr Lys Asn Val Asp Ile  
215 195 200 205  
216 Phe Asn Lys Glu Leu Leu Ile Pro Ile His Leu Glu Val His Trp  
217 210 215 220  
218 Ser Leu Ile Ser Val Asp Val Arg Arg Arg Thr Ile Thr Tyr Phe Asp  
219 225 230 235 240  
220 Ser Gln Arg Thr Leu Asn Arg Arg Cys Pro Lys His Ile Ala Lys Tyr  
221 245 250 255  
222 Leu Gln Ala Glu Ala Val Lys Lys Asp Arg Leu Asp Phe His Gln Gly

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/848,852A

DATE: 12/20/2001  
TIME: 11:52:09

Input Set : N:\CrF3\RULE60\09848852.raw  
Output Set: N:\CRF3\12202001\I848852A.raw

223 260 265 270  
224 Trp Lys Gly Tyr Phe Lys Met Asn Val Ala Arg Gln Asn Asn Asp Ser  
225 275 280 285  
226 Asp Cys Gly Ala Phe Val Leu Gln Tyr Cys Lys His Leu Ala Leu Ser  
227 290 295 300  
228 Gln Pro Phe Ser Phe Thr Gln Gln Asp Met Pro Lys Leu Arg Arg Gln  
229 305 310 315 320  
230 Ile Tyr Lys Glu Leu Cys His Cys Lys Leu Thr Val  
231 325 330

233 (2) INFORMATION FOR SEQ ID NO: 4:

235 (i) SEQUENCE CHARACTERISTICS:  
236 (A) LENGTH: 1991 base pairs  
237 (B) TYPE: nucleic acid  
238 (C) STRANDEDNESS: single  
239 (D) TOPOLOGY: linear

241 (vii) IMMEDIATE SOURCE:

242 (A) LIBRARY: BEPINOT01  
243 (B) CLONE: 2056178

245 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

247	GCCTCCCTGT	CCCCGACCC	TCTTTGATG	CCTCAGCAAG	TGAAGAGGAG	GAAGAAGAGG	60
248	AGGAGGAGGA	GGATGAAGAT	GAAGAGGAGG	AAGTGGCAGC	TTGGAGGCTG	CCCCCAAGAT	120
249	GGAGTCAGCT	GGGAACCTCC	CAGCGGCC	GCCCTTCCG	CCCCACTCAT	CGAAAAACCT	180
250	GCTCACAGCG	CCGCGGCCGA	GCCATGAGAG	CCTCCGGAT	GCTGCTCTAC	TCAAAAAGCA	240
251	CCTCGCTGAC	ATTCCACTGG	AAGCTTGGG	GGGCCACCG	GGGCCGGCG	CGGGGCCTCG	300
252	CACACCCAA	GAACCATCTT	TCACCCCAGC	AAGGGGTGC	GACGCCACAG	GTGCCATCCC	360
253	CCTGTTGTCG	TTTGACTCC	CCCCGGGGC	CACTTCCACC	CCGGCTGGGT	CTGCTAGGTG	420
254	CTCTCATGGC	TGAGGATGGG	GTGAGAGGGT	CTCCACCAGT	GCCCTCTGGG	CCCCCATGG	480
255	AGGAAGATGG	ACTCAGGTGG	ACTCCAAAGT	CTCCCTCTGGA	CCCTGACTCG	GGCCTCCTT	540
256	CATGTACTCT	GCCCAACGGT	TTTGGGGAC	AATCTGGGCC	AGAAGGGGAG	CGCACTTGGC	600
257	ACCCCTGAT	GCCAGCATCC	TCATCAGCA	TGTGTGCAGC	ATCGGGGACC	ATGTGGCCA	660
258	GGAGCTTTT	CAGGGCTCAG	ATTGGGCAT	GGCAGAAAGAG	GCAGAGAGGC	CTGGGGAGAA	720
259	AGCCGGCCAG	CACAGCCCC	TGCGAGAGGA	GCATGTGACC	TGCGTACAGA	GCATCTGGA	780
260	CGAATTCTT	CAAACGTATG	GCAGCCTCAT	ACCCCTCAGC	ACTGATGAGG	TAGTAGAGAA	840
261	GCTGGAGGAC	ATTTCAGC	AGGAGTTTC	CACCCCTCC	AGGAAGGGCC	TGGTGTGCA	900
262	GCTGATCCAG	TCTTACCAGC	GGATGCCAGG	CAATGCCATG	GTGAGGGCT	TCCGAGTGGC	960
263	TTATAAGCGG	CACGTGCTGA	CCATGGATGA	CTTGGGGACC	TTGTATGGAC	AGAACTGGCT	1020
264	CAATGACCAAG	GTGATGAACA	TGTATGGAGA	CCTGGTCATG	GACACAGTCC	CTGAAAAGGT	1080
265	GCATTCTTC	AATAGTTCT	TCTATGATAA	ACTCCGTACC	AAGGGTTATG	ATGGGGTGAA	1140
266	AAGGTGGACC	AAAAACGTGG	ACATCTCAA	TAAGGAGCTA	CTGCTAATCC	CCATCCACCT	1200
267	GGAGGTGCAT	TGGTCCCTCA	TCTCTGTGA	TGTGAGGCAG	CGCACCATCA	CCTATTTGA	1260
268	CTCGCAGCGT	ACCCCTAACCC	GCCGCTGCC	TAAGCATATT	GCCAAGTATC	TACAGGCAGA	1320
269	GGCGTAAAG	AAAGACCGAC	TGGATTTCCA	CCAGGGCTGG	AAAGGTTACT	TCAAAATGAA	1380
270	TGTGGCCAGG	CAGAATAATG	ACAGTGACTG	TGGTGTCTTT	GTGTTGCAGT	ACTGCAAGCA	1440
271	TCTGGCCCTG	TCTCAGCCAT	TCAGCTTCAC	CCAGCAGGAC	ATGCCAAAC	TTCGTCGGCA	1500
272	GATCTACAAG	GAGCTGTGTC	ACTGCAAATC	CACTGTGTGA	GCCTCGTACC	CCAGACCCCCA	1560
273	AGCCCATAAA	TGGGAAGGG	GACATGGGAG	TCCCTCCCA	AGAAACTCCA	GTCCTTTCC	1620
274	TCTCTGCCT	CTTCCCACTC	ACTTCCCTT	GGTTTTCTAT	ATTTAAATGT	TTCAATTCT	1680
275	GTATTTTTT	TTCTTGAGA	GAATACTGT	TGATTTCTGA	TGTGCAGGG	GTGGCTACAG	1740
276	AAAAGCCCT	TTCTCCTCT	GTTTGCAGGG	GAGTGTGGCC	CTGTGGCCTG	GGTGGAGCAG	1800

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/848,852A

DATE: 12/20/2001  
TIME: 11:52:10

Input Set : N:\Crf3\RULE60\09848852.raw  
Output Set: N:\CRF3\12202001\I848852A.raw

L:31 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]  
L:32 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]